



## SEQUENCE LISTING

&lt;110&gt; Li, Ming

&lt;120&gt; T-Type Calcium Channel

&lt;130&gt; 004.00191

&lt;140&gt; US 09/383,894

&lt;141&gt; 1999-08-26

&lt;150&gt; US 60/098,004

&lt;151&gt; 1998-08-26

&lt;150&gt; US 60/117,399

&lt;151&gt; 1999-01-27

&lt;160&gt; 11

&lt;170&gt; PatentIn Ver. 2.1

&lt;210&gt; 1

&lt;211&gt; 7129

&lt;212&gt; DNA

&lt;213&gt; Rattus sp.

&lt;400&gt; 1

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a!  
amt.

Phe Ile Phe Ser Ile Leu Gly Met His Leu Phe Gly Cys Lys Phe Ala  
 930 935 940  
 Ser Glu Arg Asp Gly Asp Thr Leu Pro Asp Arg Lys Asn Phe Asp Ser  
 945 950 955 960  
 Leu Leu Trp Ala Ile Val Thr Val Phe Gln Ile Leu Thr Gln Glu Asp  
 965 970 975  
 Trp Asn Lys Val Leu Tyr Asn Gly Met Ala Ser Thr Ser Ser Trp Ala  
 980 985 990  
 Ala Leu Tyr Phe Ile Ala Leu Met Thr Phe Gly Asn Tyr Val Leu Phe  
 995 1000 1005  
 Asn Leu Leu Val Ala Ile Leu Val Glu Gly Phe Gln Ala Glu Glu Ile  
 1010 1015 1020  
 Gly Lys Arg Glu Asp Ala Ser Gly Gln Leu Ser Cys Ile Gln Leu Pro  
 1025 1030 1035 1040  
 Val Asn Ser Gln Gly Gly Asp Ala Thr Lys Ser Glu Ser Glu Pro Asp  
 1045 1050 1055  
 Phe Phe Ser Pro Ser Val Asp Gly Asp Gly Asp Arg Lys Lys Arg Leu  
 1060 1065 1070  
 Ala Leu Val Ala Leu Gly Glu His Ala Glu Leu Arg Lys Ser Leu Leu  
 1075 1080 1085  
 Pro Pro Leu Ile Ile His Thr Ala Ala Thr Pro Met Ser Leu Pro Lys  
 1090 1095 1100  
 Ser Ser Ser Thr Gly Val Gly Glu Ala Leu Gly Ser Gly Ser Arg Arg  
 1105 1110 1115 1120  
 Thr Ser Ser Ser Gly Ser Ala Glu Pro Gly Ala Ala His His Glu Met  
 1125 1130 1135  
 Lys Ser Pro Pro Ser Ala Arg Ser Ser Pro His Ser Pro Trp Ser Ala  
 1140 1145 1150  
 Ala Ser Ser Trp Thr Ser Arg Arg Ser Ser Arg Asn Ser Leu Gly Arg  
 1155 1160 1165  
 Ala Pro Ser Leu Lys Arg Arg Ser Pro Ser Gly Glu Arg Arg Ser Leu  
 1170 1175 1180

A!  
 cont.

Leu Ser Gly Glu Gly Gln Glu Ser Gln Asp Glu Glu Glu Ser Ser Glu  
 1185 1190 1195 1200  
 Glu Asp Arg Ala Ser Pro Ala Gly Ser Asp His Arg His Arg Gly Ser  
 1205 1210 1215  
 Leu Glu Arg Glu Ala Lys Ser Ser Phe Asp Leu Pro Asp Thr Leu Gln  
 1220 1225 1230  
 Val Pro Gly Leu His Arg Thr Ala Ser Gly Arg Ser Ser Ala Ser Glu  
 1235 1240 1245  
 His Gln Asp Cys Asn Gly Lys Ser Ala Ser Gly Arg Leu Ala Arg Thr  
 1250 1255 1260  
 Leu Arg Thr Asp Asp Pro Gln Leu Asp Gly Asp Asp Asp Asn Asp Glu  
 1265 1270 1275 1280  
 Gly Asn Leu Ser Lys Gly Glu Arg Ile Gln Ala Trp Val Arg Ser Arg  
 1285 1290 1295  
 Leu Pro Ala Cys Cys Arg Glu Arg Asp Ser Trp Ser Ala Tyr Ile Phe  
 1300 1305 1310  
 Pro Pro Gln Ser Arg Phe Arg Leu Leu Cys His Arg Ile Ile Thr His  
 1315 1320 1325  
 Lys Met Phe Asp His Val Val Leu Val Ile Ile Phe Leu Asn Cys Ile  
 1330 1335 1340  
 Thr Ile Ala Met Glu Arg Pro Lys Ile Asp Pro His Ser Ala Glu Arg  
 1345 1350 1355 1360  
 Ile Phe Leu Thr Leu Ser Asn Tyr Ile Phe Thr Ala Val Phe Leu Ala  
 1365 1370 1375  
 Glu Met Thr Val Lys Val Val Ala Leu Gly Trp Cys Phe Gly Glu Gln  
 1380 1385 1390  
 Ala Tyr Leu Arg Ser Ser Trp Asn Val Leu Asp Gly Leu Leu Val Leu  
 1395 1400 1405  
 Ile Ser Val Ile Asp Ile Leu Val Ser Met Val Ser Asp Ser Gly Thr  
 1410 1415 1420  
 Lys Ile Leu Gly Met Leu Arg Val Leu Arg Leu Leu Arg Thr Leu Arg  
 1425 1430 1435 1440

a!  
 cont.

Pro Leu Arg Val Ile Ser Arg Ala Gln Gly Leu Lys Leu Val Val Glu  
 1445 1450 1455  
 Thr Leu Met Ser Ser Leu Lys Pro Ile Gly Asn Ile Val Val Ile Cys  
 1460 1465 1470  
 Cys Ala Phe Phe Ile Ile Phe Gly Ile Leu Gly Val Gln Leu Phe Lys  
 1475 1480 1485  
 Gly Lys Phe Phe Val Cys Gln Gly Glu Asp Thr Arg Asn Ile Thr Asn  
 1490 1495 1500  
 Lys Ser Asp Cys Ala Glu Ala Ser Tyr Arg Trp Val Arg His Lys Tyr  
 1505 1510 1515 1520  
 Asn Phe Asp Asn Leu Gly Gln Ala Leu Met Ser Leu Phe Val Leu Ala  
 1525 1530 1535  
 Ser Lys Asp Gly Trp Val Asp Ile Met Tyr Asp Gly Leu Asp Ala Val  
 1540 1545 1550  
 Gly Val Asp Gln Gln Pro Ile Met Asn His Asn Pro Trp Met Leu Leu  
 1555 1560 1565  
 Tyr Phe Ile Ser Phe Leu Leu Ile Val Ala Phe Phe Val Leu Asn Met  
 1570 1575 1580  
 Phe Val Gly Val Val Val Glu Asn Phe His Lys Cys Arg Gln His Gln  
 1585 1590 1595 1600  
 Glu Glu Glu Glu Ala Arg Arg Arg Glu Glu Lys Arg Leu Arg Arg Leu  
 1605 1610 1615  
 Glu Lys Lys Arg Arg Asn Leu Met Leu Asp Asp Val Ile Ala Ser Gly  
 1620 1625 1630  
 Ser Ser Ala Ser Ala Ala Ser Glu Ala Gln Cys Lys Pro Tyr Tyr Ser  
 1635 1640 1645  
 Asp Tyr Ser Arg Phe Arg Leu Leu Val His His Leu Cys Thr Ser His  
 1650 1655 1660  
 Tyr Leu Asp Leu Phe Ile Thr Gly Val Ile Gly Leu Asn Val Val Thr  
 1665 1670 1675 1680  
 Met Ala Met Glu His Tyr Gln Gln Pro Gln Ile Leu Asp Glu Ala Leu  
 1685 1690 1695

Lys Ile Cys Asn Tyr Ile Phe Thr Val Ile Phe Val Phe Glu Ser Val  
 1700 1705 1710  
 Phe Lys Leu Val Ala Phe Gly Phe Arg Arg Phe Phe Gln Asp Arg Trp  
 1715 1720 1725  
 Asn Gln Leu Asp Leu Ala Ile Val Leu Leu Ser Ile Met Gly Ile Thr  
 1730 1735 1740  
 Leu Glu Glu Ile Glu Val Asn Ala Ser Leu Pro Ile Asn Pro Thr Ile  
 1745 1750 1755 1760  
 Ile Arg Ile Met Arg Val Leu Arg Ile Ala Arg Val Leu Lys Leu Leu  
 1765 1770 1775  
 Lys Met Ala Val Gly Met Arg Ala Leu Leu Asp Thr Val Met Gln Ala  
 1780 1785 1790  
 Leu Pro Gln Val Gly Asn Leu Gly Leu Leu Phe Met Leu Leu Phe Phe  
 1795 1800 1805  
 Ile Phe Ala Ala Leu Gly Val Glu Leu Phe Gly Asp Leu Glu Cys Asp  
 1810 1815 1820  
 Glu Thr His Pro Cys Glu Gly Leu Gly Arg His Ala Thr Phe Arg Asn  
 1825 1830 1835 1840  
 Phe Gly Met Ala Phe Leu Thr Leu Phe Arg Val Ser Thr Gly Asp Asn  
 1845 1850 1855  
 Trp Asn Gly Ile Met Lys Asp Thr Leu Arg Asp Cys Asp Gln Glu Ser  
 1860 1865 1870  
 Thr Cys Tyr Asn Thr Val Ile Ser Pro Ile Tyr Phe Val Ser Phe Val  
 1875 1880 1885  
 Leu Thr Ala Gln Phe Val Leu Val Asn Val Val Ile Ala Val Leu Met  
 1890 1895 1900  
 Lys His Leu Glu Glu Ser Asn Lys Glu Ala Lys Glu Glu Ala Glu Leu  
 1905 1910 1915 1920  
 Glu Ala Glu Leu Glu Leu Glu Met Lys Thr Leu Ser Pro Gln Pro His  
 1925 1930 1935  
 Ser Pro Leu Gly Ser Pro Phe Leu Trp Pro Gly Val Glu Gly Val Asn  
 1940 1945 1950

Ser Pro Asp Ser Pro Lys Pro Gly Ala Pro His Thr Thr Ala His Ile  
 1955 1960 1965  
 Gly Ala Ala Ser Gly Phe Ser Leu Glu His Pro Thr Met Val Pro His  
 1970 1975 1980  
 Pro Glu Glu Val Pro Val Pro Leu Gly Pro Asp Leu Leu Thr Val Arg  
 1985 1990 1995 2000  
 Lys Ser Gly Val Ser Arg Thr His Ser Leu Pro Asn Asp Ser Tyr Met  
 2005 2010 2015  
 Cys Arg Asn Gly Ser Thr Ala Glu Arg Ser Leu Gly His Arg Gly Trp  
 2020 2025 2030  
 Gly Leu Pro Lys Ala Gln Ser Gly Ser Ile Leu Ser Val His Ser Gln  
 2035 2040 2045  
 Pro Ala Asp Thr Ser Cys Ile Leu Gln Leu Pro Lys Asp Val His Tyr  
 2050 2055 2060  
 Leu Leu Gln Pro His Gly Ala Pro Thr Trp Gly Ala Ile Pro Lys Leu  
 2065 2070 2075 2080  
 Pro Pro Pro Gly Arg Ser Pro Leu Ala Gln Arg Pro Leu Arg Arg Gln  
 2085 2090 2095  
 Ala Ala Ile Arg Thr Asp Ser Leu Asp Val Gln Gly Leu Gly Ser Arg  
 2100 2105 2110  
 Glu Asp Leu Leu Ser Glu Val Ser Gly Pro Ser Cys Pro Leu Thr Arg  
 2115 2120 2125  
 Ser Ser Ser Phe Trp Gly Gly Ser Ser Ile Gln Val Gln Gln Arg Ser  
 2130 2135 2140  
 Gly Ile Gln Ser Lys Val Ser Lys His Ile Arg Leu Pro Ala Pro Cys  
 2145 2150 2155 2160  
 Pro Gly Leu Glu Pro Ser Trp Ala Lys Asp Pro Pro Glu Thr Arg Ser  
 2165 2170 2175  
 Ser Leu Glu Leu Asp Thr Glu Leu Ser Trp Ile Ser Gly Asp Leu Leu  
 2180 2185 2190  
 Pro Ser Ser Gln Glu Glu Pro Leu Ser Pro Arg Asp Leu Lys Lys Cys  
 2195 2200 2205

a'  
 cont.



Tyr Ser Val Glu Thr Gln Ser Cys Arg Arg Arg Pro Gly Ser Trp Leu  
 2210 2215 2220

Asp Glu Gln Arg Arg His Ser Ile Ala Val Ser Cys Leu Asp Ser Gly  
 2225 2230 2235 2240

Ser Gln Pro Arg Leu Cys Pro Ser Pro Ser Ser Leu Gly Gly Gln Pro  
 2245 2250 2255

Leu Gly Gly Pro Gly Ser Arg Pro Lys Lys Lys Leu Ser Pro Pro Ser  
 2260 2265 2270

Ile Ser Ile Asp Pro Pro Glu Ser Gln Gly Ser Arg Pro Pro Cys Ser  
 2275 2280 2285

Pro Gly Val Cys Leu Arg Arg Arg Ala Pro Ala Ser Asp Ser Lys Asp  
 2290 2295 2300

Pro Ser Val Ser Ser Pro Leu Asp Ser Thr Ala Ala Ser Pro Ser Pro  
 2305 2310 2315 2320

Lys Lys Asp Thr Leu Ser Leu Ser Gly Leu Ser Ser Asp Pro Thr Asp  
 2325 2330 2335

Met Asp Pro Val Leu Pro Thr Leu Pro His His Leu Ser Pro Pro Gly  
 2340 2345 2350

Ala Asp Pro Ser Ser Ala Ser Trp Ala Ala Phe Leu Lys Ser Pro Thr  
 2355 2360 2365

Ala Ala Ser Ser His Glu Ala Pro His Leu Pro Ser Ser Val Ala Gly  
 2370 2375 2380

Gly Asp Asp Glu Gln Asn Phe Arg Arg Val Asp Leu Lys Arg Thr Gln  
 2385 2390 2395 2400

Pro Trp Ser Pro Cys Leu Arg Glu Glu Gly Lys Gly Glu Ser Pro Val  
 2405 2410 2415

Trp Pro Arg Leu Pro Thr Pro Gly Ala  
 2420 2425

<210> 5

<211> 7

<212> PRT

<213> Rattus sp.

<400> 5  
Ser Lys Glu Lys Gln Met Ala  
1 5

<210> 6  
<211> 17  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: Primer  
Sequence

<400> 6  
tngchatgga gmgnc cy

17

<210> 7  
<211> 18  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: Primer  
Sequence

<400> 7  
cttbcccttg aasarttg

18

<210> 8  
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<212> DNA  
<213> Artificial Sequence

<220>  
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Sequence

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ccgctgtcgg agaccatgga gacc

24

<210> 9  
<211> 25  
<212> DNA  
<213> Artificial Sequence

Al  
Cmt

<220>

<223> Description of Artificial Sequence: Primer  
Sequence

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25

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<211> 18

<212> DNA

<213> Artificial Sequence

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Sequence

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18

<210> 11

<211> 18

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer  
Sequence

<400> 11

ctgtggcgat ggtcactg

18